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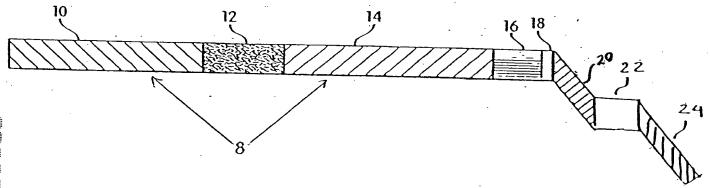
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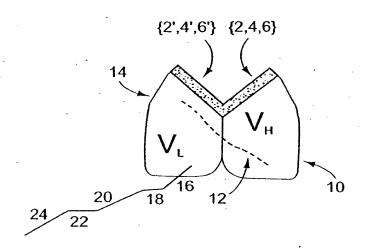
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SINGLE CHAIN BINDING POLYPEPTIDE.

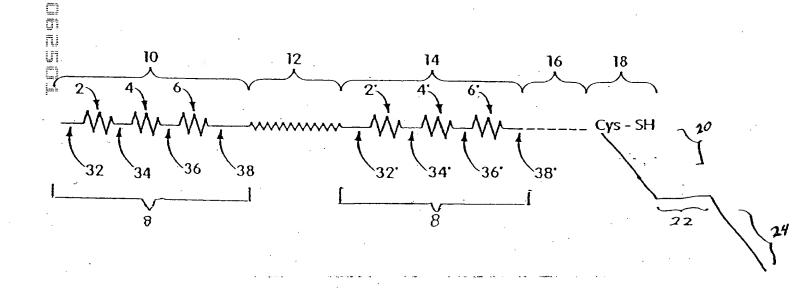


(a) Extended Polypeptide



(b) Folded Protein

SINGLE CHAIN
BINDING POLYPEPTIDE SHOWING
LOCATIONS OF COMPLEMENTARITY
DETERMINING REGIONS, POLYPEPTIDE
SPACER REGIONS, AND EFFECTOR REGIONS



C6.5 sFv AMINO ACID SEQUENCE

(N-terminus to C-terminus)

-QVQLLQSGAE LKKPGESLKI SCKGSGYSFT SYWIAWVRQM PGKGLEYMGL IYPGDSDTKY SPSFQGQVTI SVDKSVSTAY LQWSSLKPSD SAVYFCARHD VGYCSSSNCA KWPEYFQHWG QGTLVTVSSG GGGSGGGGSG GGGSQSVLTQ PPSVSAAPGQ KVTISCSGSS SNIGNNYVSW YQQLPGTAPK LLIYGHTNRP AGVPDRFSGS KSGTSASLAI SGFRSEDEAD YYCAAWDDSL SGWVFGGGTK LTVLG

FIGURE 4

C6.5 sFv NUCLEOTIDE SEQUENCE

C6ML3-9 sFv' AMINO ACID SEQUENCE

(N-terminus to C-terminus)

-QVQLVQSGAE VKKPGESLKI SCKGSGYSFT SYWIAWVRQM PGKGLEYMGL IYPGDSDTKY SPSFQGQVTI SVDKSVSTAY LQWSSLKPSD SAVYFCARHD VGYCSSSNCA KWPEYFQHWG QGTLVTVSSG GGGSGGGGSG GGGSQSVLTQ PPSVSAAPGQ KVTISCSGSS SNIGNNYVSW YQQLPGTAPK LLIYDHTNRP AGVPDRFSGS KSGTSASLAI SGFRSEDEAD YYCASWDYTL SGWVFGGGTK LTVLGAAAHH HHHHGGGGC-

FIGURE 6

C6ML3-9 sF_V' NUCLEOTIDE SEQUENCE

5' caggtgcagctggtgcagtctggggcagaggtgaaaaagcccggggagtctctgaagatctcctgtaagggttctggata cagctttaccagctactggatcgcctggtgcgccagatgcccgggaaaggcctggagtacatggggctcatctatcctg gtgactctgacaccaaatacagcccgtccttccaaggccaggtcaccatctcagtcgacaagtccgtcagcactgcctac ttgcaatggagcagtctgaagccctcggacagcgccgtgtatttttgtgcgagacatgacgtgggatattgcagtagttc caactgcgcaaagtggcctgaatacttccagcattggggccagggcaccctggtcaccgtctcctcaggtggaggcggtt caggcggaggtggctctggcggtggcggatcgcagtctgtgttgacgcagcgccctcagtgtctgcggccccaggacag aaggtcaccatctcctgctctggaagcagctccaacattgggaataattatgtatcctggtaccagcagctccaagaac agccccaaaactcctcatctatgatcacaccaatcggccgaagggtccctgaccgattctctggctccaagtctggcacctcagctcctggcaccatctcctggcaccatcaccatctcggcccgaggatgaggtgattattactgtgcctcctgggactacaccctc tcgggctgggtgttcggcggaggaaccaagctgaccgtcctaggtggggcgcacaccatcaccatcacggtggtgg cggctgc3'

C6ML3-9sFv'-L1-KDEL AMINO ACID SEQUENCE

(N-terminus to C-terminus)

-QVQLVQSGAE VKKPGESLKI SCKGSGYSFT SYWIAWVRQM PGKGLEYMGL IYPGDSDTKY SPSFQGQVTI SVDKSVSTAY LQWSSLKPSD SAVYFCARHD VGYCSSSNCA KWPEYFQHWG QGTLVTVSSG GGGSGGGGSG GGGSQSVLTQ PPSVSAAPGQ KVTISCSGSS SNIGNNYVSW YQQLPGTAPK LLIYDHTNRP AGVPDRFSGS KSGTSASLAI SGFRSEDEAD YYCASWDYTL SGWVFGGGTK LTVLGAAAHH HHHHGGGGCL ESSSSGSEKD EL-

FIGURE 8

C6ML3-9 sFv'-L1-KDEL NUCLEOTIDE SEQUENCE

C6ML3-9 sFv'-L2-KDEL AMINO ACID SEQUENCE

(N-terminus to C-terminus)

-QVQLVQSGAE VKKPGESLKI SCKGSGYSFT SYWIAWVRQM PGKGLEYMGL IYPGDSDTKY SPSFQGQVTI SVDKSVSTAY LQWSSLKPSD SAVYFCARHD VGYCSSSNCA KWPEYFQHWG QGTLVTVSSG GGGSGGGGSG GGGSQSVLTQ PPSVSAAPGQ KVTISCSGSS SNIGNNYVSW YQQLPGTAPK LLIYDHTNRP AGVPDRFSGS KSGTSASLAI SGFRSEDEAD YYCASWDYTL SGWVFGGGTK LTVLGAAAHH HHHHGGGGCL ESSSSGSSSS GSEKDEL-

FIGURE 10

C6ML3-9sFv'-L2-KDEL NUCLEOTIDE SEQUENCE

FIGURE 11

C6ML3-9 sFv'-L2-H14 AMINO ACID SEQUENCE

(N-terminus to C-terminus)

-QVQLVQSGAE VKKPGESLKI SCKGSGYSFT SYWIAWVRQM PGKGLEYMGL IYPGDSDTKY SPSFQGQVTI SVDKSVSTAY LQWSSLKPSD SAVYFCARHD VGYCSSSNCA KWPEYFQHWG QGTLVTVSSG GGGSGGGGSG GGGSQSVLTQ PPSVSAAPGQ KVTISCSGSS SNIGNNYVSW YQQLPGTAPK LLIYDHTNRP AGVPDRFSGS KSGTSASLAI SGFRSEDEAD YYCASWDYTL SGWVFGGGTK LTVLGAAAHH HHHHGGGGCL ESSSSGSSSS GSKKSAKKTP KKAKKP-

FIGURE 12

C6ML3-9 sFv' -L2-H14 NUCLEOTIDE SEOUENCE

5' caggtgcagctggtgcagtctggggcagaggtgaaaaagcccggggagtctctgaagatctcctgtaagggttctggata cagctttaccagctactggatcgcctgggtgcgccagatgcccgggaaaggcctggagtacatggggctcatctatcctg gtgactctgacaccaaatacagccgtccttccaaggccaggtcaccatctcagtcgacaagtccgtcagcactgcctac ttgcaatggagcagtctgaagccctcggacagcccgtgtatttttgtgcgagacatgacgtgggatattgcagtagttc caactgcgcaaagtggcctgaatacttccagcattggggccagggcaccctggtcaccgtctcctcaggtggaggcggtt caggcggaggtggctctggcggatcgcagttgtgtgagccggccccaggacag aaggtcaccatctcctggtaggagcagctccaacattgggaataattatgtatcctggtaccagcagctcccaggaac agccccaaactcctctgtctggaagcagctccaacattgggaataattatgtatcctggtaccagcagctccaaggaac agccccaaactcctcatctatgatcacaccaatcggcccgcaggggtccctgaccgattctctggctccaagtctggca cctcagcctcctggccatcagtgggttccggtccgaggatgaggctgattattactgtgcctcctgggactacaccctc tcgggctgggtgttcgggcggaggaaccaagctgccgacgacgaccaccatcatcaccatcacggtggtgg cggctgc ctcgagtcta gcagctccgg ttcctctagc tctggatcca agaaaagcgc gaaaaagacc ccgaagaaag cgaagaaacc g 3'

C6ML3-9sFv'-L2-nls AMINO ACID SEQUENCE

(N-terminus to C-terminus)

-QVQLVQSGAE VKKPGESLKI SCKGSGYSFT SYWIAWVRQM PGKGLEYMGL IYPGDSDTKY SPSFQGQVTI SVDKSVSTAY LQWSSLKPSD SAVYFCARHD VGYCSSSNCA KWPEYFQHWG QGTLVTVSSG GGGSGGGGSG GGGSQSVLTQ PPSVSAAPGQ KVTISCSGSS SNIGNNYVSW YQQLPGTAPK LLIYDHTNRP AGVPDRFSGS KSGTSASLAI SGFRSEDEAD YYCASWDYTL SGWVFGGGTK LTVLGAAAHH HHHHGGGGCL ESSSSGSSSS GSTPPKKKRK V

FIGURE 14

C6ML3-9 sFv'-L2-nls NUCLEOTIDE SEQUENCE

5' caggtgeagctggtgcagtctggggcagaggtgaaaaagccgggggagtctctgaagatctcctgtaagggttctggata cagctttaccagctactggatcgcctgggtgcgccagatgccgggaaaagcctggagtacatggggctcatctatcctg gtgactctgacaccaaatacagcccgtcettccaaggccaggtcaccatctcagtcgacaagtccgtcagcactgcctac ttgcaatggagcagtctgaagccctcggacagcgccgtgtatttttgtgcgagacatgacgtgggatattgcagtagttc caactgcgcaaagtggcctgaatacttccagcattggggccagggcaccctggtcaccgtctcctcaggtggaggcggtt caggcggaggtggctctggcggtggcggatcgcagtctgtgttgacgcagcgccctcagtgtctgcggccccaggacag aaggtcaccatctcctgctctggaagcagctccaacattgggaataattatgtatcctggtaccagcagctccaagtctggca agccccaaactcctcatctatgatcacaccaatcggccgcaggggtccctgaccgattctctggctccaagtctggca cctcagcctccctggccatcagtgggttccggtccgaggatgaggctgattattactgtgcctcctgggactacaccctc tcgggctggtgttcggcggaggaaccaagctgccaggtgggggggcgccacaccatcaccaccatcacggtggtgg cggctgc ctcgagtcta gcagctccgg ttcctcagc tctggatca ctccgccgaa aāāgaaacgt aaagtg 3'

Figure 15. C6ML3-9 sFv' and its salmon protamine conjugate binds specifically to the erbB-2 positive ovarian cancer cells

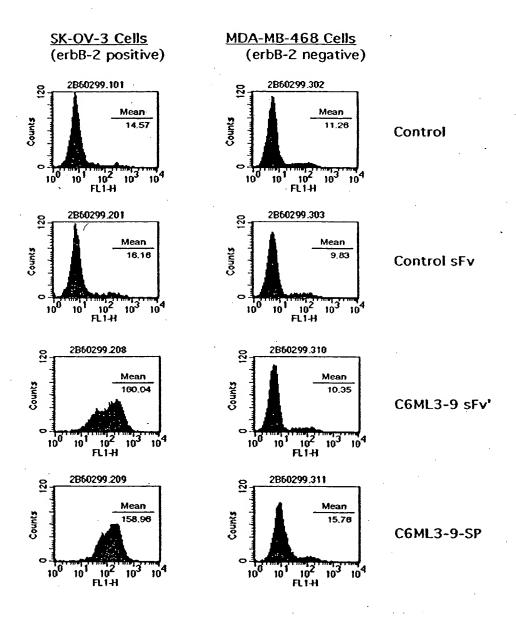


Figure 16. FACS Analysis of the erbB-2 Binding Activities of Bacterially Expressed C6ML3-9 sFv' and its Derivatives

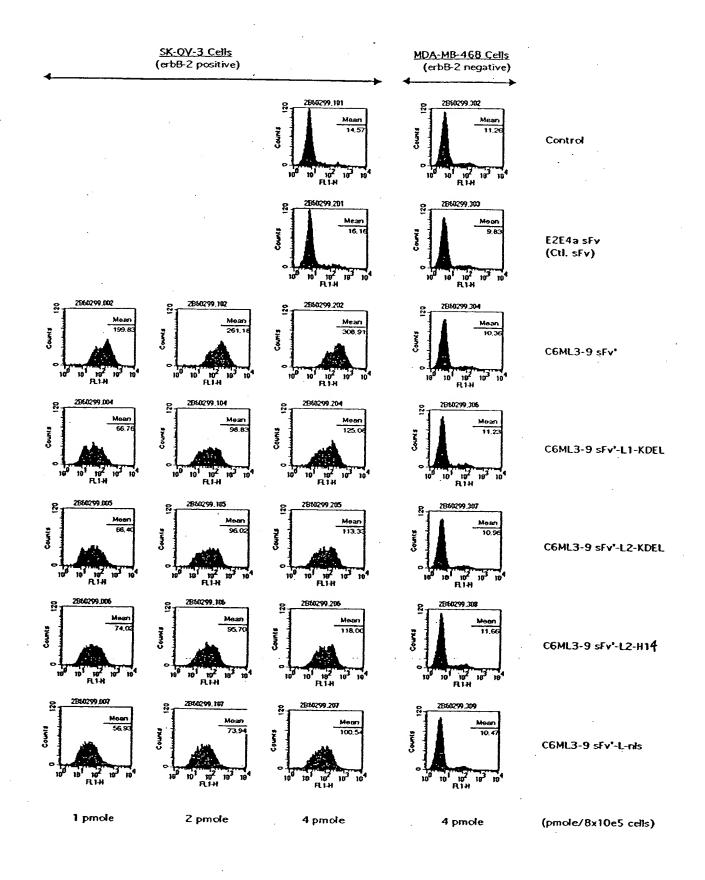
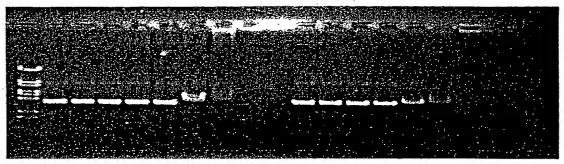


Figure 17. Gel Shift Analysis of the C6.5-SP-DNA and C6ML3-9-SP-DNA Complex

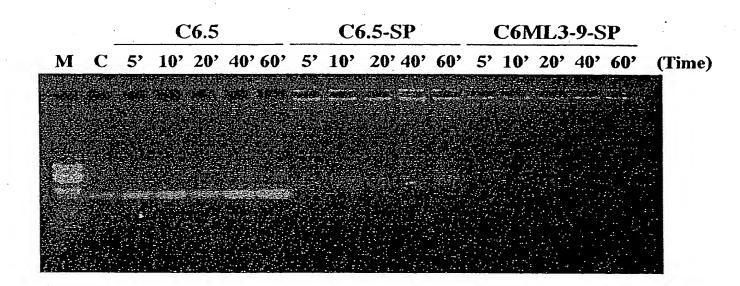
M 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17



- M. DNA marker λ DNA BstEll digest
- 1. 200 ng pGL3 DNA
- 2. 200 ng pGL3 DNA + 1.45 pmol C6.5
- 3. 200 ng pGL3 DNA + 2.90 pmol C6.5
- 4. 200 ng pGL3 DNA + 5.80 pmol C6.5
- 5. 200 ng pGL3 DNA + 11.6 pmol C6.5
- 6. 200 ng pGL3 DNA + 1.45 pmol C6.5-SP
- 7. 200 ng pGL3 DNA + 2.90 pmol C6.5-SP
- 8. 200 ng pGL3 DNA + 5.80 pmol C6.5-SP
- 9. 200 ng pGL3 DNA + 11.6 pmol C6.5-SP
- 10. 200 ng pGL3 DNA + 1.45 pmol C6ML3-9
- 11. 200 ng pGL3 DNA + 2.90 pmol C6ML3-9
- 12. 200 ng pGL3 DNA + 5.80 pmol C6ML3-9
- 13. 200 ng pGL3 DNA + 11.6 pmol C6ML3-9
- 14. 200 ng pGL3 DNA + 1.45 pmol C6ML3-9-SP
- 15. 200 ng pGL3 DNA + 2.90 pmol C6ML3-9-SP
- 16. 200 ng pGL3 DNA + 5.80 pmol C6ML3-9-SP
- 17. 200 ng pGL3 DNA + 11.6 pmol C6ML3-9-SP

^{*0.8%} agarose gel in 1xTAE, 150v, RT, ~lhr, EtBr staining overnight

Figure 18. Kinetic Study of the C6.5-SP-DNA and C6ML3-9-SP-DNA Complex Formation



- M. DNA marker λ DNA BstEII digest
- C. 200 ng pGL3 DNA alone
- * The rest of the lanes 200 ng pGL3 DNA incubated with 5.8 pmol proteins as indicated above each line, on ice, for different period of time. Electrophoresis condition same as Figure 3.

Figure 19. The C6ML3-9-SP conjugate protein mediates specific luciferase gene delivery to erbB-2 positive cancer cells

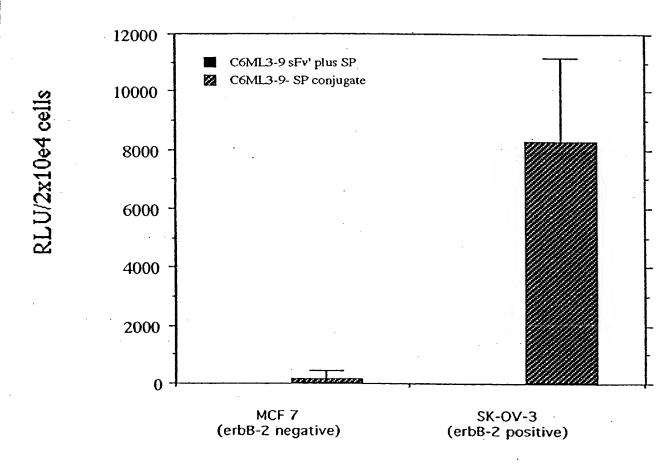


Figure 20. Chloroquine-dependent C6ML3-9-SP-mediated Gene Delivery

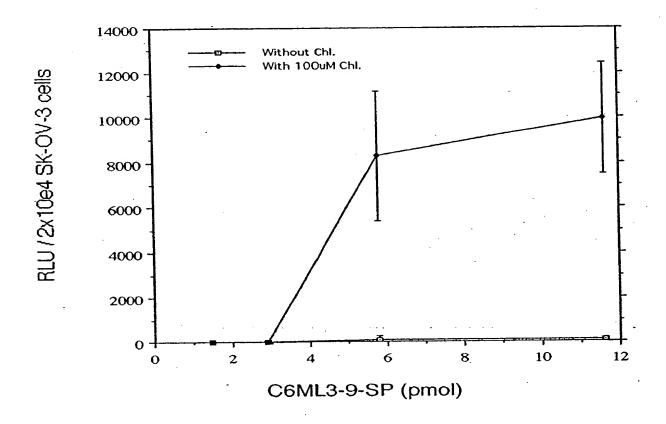
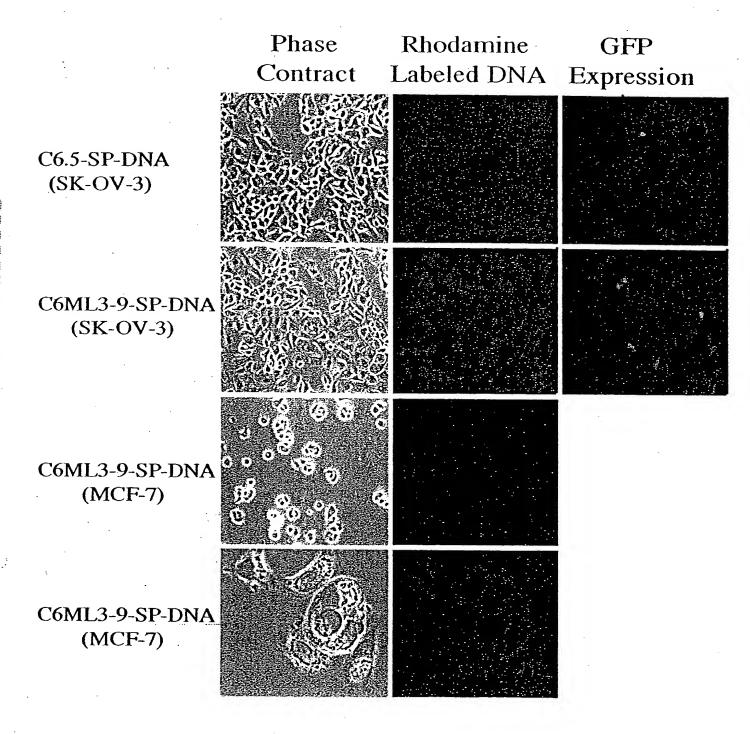


Figure-21. Fluorescent microscopy of 6.5-SP and C6ML3-9 -SP-mediated gene transfer of pGeneGrip Rhodamine/GFP plasmids with SK-OV-3 and MCF-7



THE EFFECT OF CHLOROQUINE ON'3T3-HER2 TRANSFECTION MEDIATED BY C6ML3-9sFv'-SALMON PROTAMINE

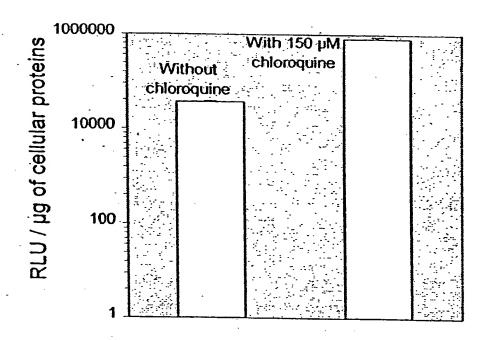
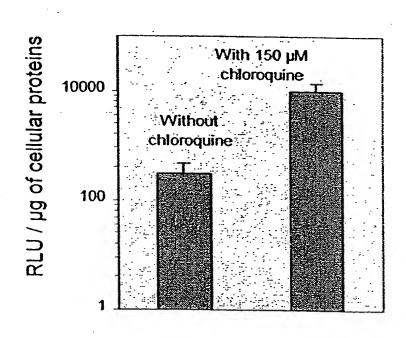
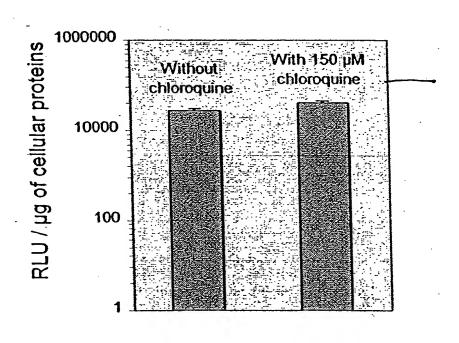


FIGURE 23

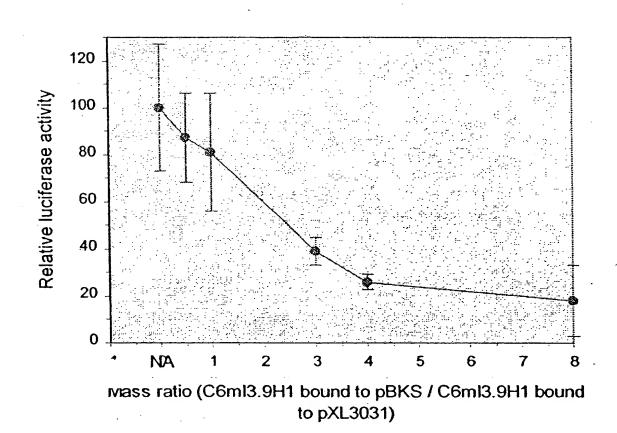
THE EFFECT OF CHLOROQUINE ON 3T3-HER2 TRANSFECTION MEDIATED BY C6ML3-9sFv'#2-P1



THE EFFECT OF CHLOROQUINE ON 3T3-HER2 TRANSFECTION MEDIATED BY C6ML3-9sFv'#2-H1



THE EFFECT OF C6ML3-9sFv'-H1-pBks ON 3T3-HER2 TRANSFECTION MEDIATED BY C6ML3-9sFv'-H1



THE EFFECT OF THE DNA TO C6ML3-9sFv'-H1 RATIO ON 3T3-HER2 TRANSFECTION EFFICIENCY

